```
logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        MLMHHYKQKAQAGVGLLEVLVALILLAIGVLGYVALQLRAMDASSEALSKSQAILVMRGLAENIRTNSTQA..SQYPTFV    78
NTDB id 1094081 ACMHYC RS02180 WP 414659618.1  ......MKIKQQGVGLVEVLVALVLLAISVLGFAALQLRAVDSSLEANKQVIALTIARDLAEKMRANPQATRAGKYKVAA    74
NTDB id 1003 ACIAD RS15025 WP 004923832.1       ....MSKMRAQRGVGMVEVLVALLLLAIGVLGFTALQLRAVSASMEASNQVLALNLARDLTERMRANPQATQCKLYHTEG    76
consensus                                              * **! !!!**!!!!!!*!!!!*!!!* !!!!!!***!*!!  ***!* **!*!*!**!*!****    ! * *

logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        RSYSNYTSDTPAPTSCFNSLCT.ASQLAQFDAYQAARNANQLGMRITMSNCPGVTNTM.VQQRQCLFVFWGKTAPVITTN   156
NTDB id 1094081 ACMHYC RS02180 WP 414659618.1  VEVP.KTSVS........NVADKPEKIAIQDLQNAALQAKDSGMNLAVEQCPTGN......NRQCIYVAWDETNLAGND.   138
NTDB id 1003 ACIAD RS15025 WP 004923832.1       ALAK.DTKAYTKASNCSGSVSTQATAQANSDLLFIYNQANTLGMAVIINTCPNATNAGNNLGRQCVYIAWDKTAQNFTN.   154
consensus                                              !* *   * *  ** * ** *!* !* *** *!* *!! * * *!! ***      *!!!****!**!*   *

logo
NTDB id 1067 ABD1 RS15745 WP 002194578.1        GTNTSVDVSSCM.SNNGVYVNNSTCLMMEAY   186
NTDB id 1094081 ACMHYC RS02180 WP 414659618.1  ......GIASCIKD..GAYVSGSHCLFLEAY   161
NTDB id 1003 ACIAD RS15025 WP 004923832.1       .......SDTCYKSGGSSYERNTTCLFLEAY   178
consensus                                              * *!***  * !* ***!!**!!!
```

X non conserved

X similar

X $\geq$50% conserved